

SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name _____ Examiner # _____ Date: _____
 Att Unit _____ Phone Number 30 _____ Serial Number _____
 Mail Box and Bldg Room Location _____ Results Format Preferred (circle): PAPER DISK E-MAIL

If more than one search is submitted, please prioritize searches in order of need.

Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the electron species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc., if known. Please attach a copy of the cover sheet, pertinent claims, and abstract.

Title of Invention: _____

Inventors (please provide full names): _____

Earliest Priority Filing Date: _____

**For Sequence Searches Only* Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number.*

BEST AVAILABLE COPY

STAFF USE ONLY

Searcher: P. Schreiber
 Searcher Phone #: 308-4292
 Searcher Location: CMC C103
 Date: 10/8
 Searcher Fee & Fee Waiver: 13
 Searcher Prep Time: _____
 Searcher Time: 16

Type of Search

NA Sequence (P) _____
 AA Sequence (P) 2
 Structure (P) _____
 Bibliographic _____
 Litigation _____
 Full text _____
 Patent Family _____
 Other _____

Vendors and cost where applicable

STN _____
 Dialog _____
 Quest _____
 JACS Net _____
 Sequences Systems CompuGen 606
 WWW Internet ✓
 Other _____

Schreiber, David

105875

From: Ramirez, Delia
Sent: Wednesday, October 08, 2003 3:01 PM
To: Schreiber, David
Subject: case 09/856679

Hi David,

I was wondering if you could do the following alignment. I need baa04770 gi474982 April 27, 1994 against seq id 2 and the corresponding DNA against seq id 31.

Thank you,

Delia M. Ramirez, Ph.D.
Patent Examiner
Recombinant Enzymes-Art Unit 1652
USPTO
1911 S. Clark Street, Crystal Mall 1, 10D06, Mail room 10D01
Arlington, VA 22202
(703) 306-0288
delia.ramirez@uspto.gov

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Scientific and Technical Information Center

Requester's Full Name _____ Examiner # _____ Date: _____
 An Unit _____ Phone Number 30 _____ Serial Number _____
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Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc., if known. Please attach a copy of the cover sheet, pertinent claims, and abstract.

Title of Invention: _____

Inventors (please provide full names): _____

Earliest Priority Filing Date: _____

**For Sequence Searches Only* Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number*

BEST AVAILABLE COPY

STAFF USE ONLY

Searcher: P. Schwab
 Searcher Phone #: 308-4292
 Searcher Location: CM 6A03
 Date: 10/8
 Searcher Prior Filing Time: 12
 Date: 18

Type of Search

NA Sequence #: 2
 AA Sequence #: 2
 Structure #: _____
 Bibliographic: _____
 Litigation: _____
 Full-text: _____
 Patent Family: _____
 Other: _____

Vendors and cost where applicable

STN: _____
 Dialog: _____
 Quest: _____
 Compaq: _____
 WPA: _____
 Other: _____

Schreiber, David

5874

From: Ramirez, Delia
Sent: Wednesday, October 08, 2003 5:29 PM
To: Schreiber, David
Subject: case 09/856,679

Hi David,

I would like to request the following alignments:

AC004241 gi 2920325 Feb 1998 against seq id 31
AC004241 gi 3282154 July 1998 against seq id 31
AC004241 gi 3108007 May 2 1998 against seq id 31
AC004241 gi 3046263 Apr 13 1998 against seq id 31
AC004241 gi 3023023 Apr 4 1998 against seq id 31
AC004241 gi 3004524 Apr 1 1998 against seq id 31
AC004241 gi 2995474 Mar 28 1998 against seq id 31
AC004241 gi 2980958 Mar 21 1998 against seq id 31
AC004241 gi 2960506 Mar 14 1998 against seq id 31
AC004241 gi 2943840 Mar 7 1998 against seq id 31

Thank you,

Delia M. Ramirez, Ph.D.
Patent Examiner
Recombinant Enzymes-Art Unit 1652
USPTO
1911 S. Clark Street, Crystal Mall 1, 10D06, Mail room 10D01
Arlington, VA 22202
(703) 306-0288
delia.ramirez@uspto.gov

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OM protein - protein search, using sw model

Run on: October 8, 2003, 15:50:09 ; Search time 0.001 Seconds
(without alignments)
364.026 Million cell updates/sec

Title: us-09-856-679-2

Perfect score: 1759
Sequence: 1 MALAQEDGWTGQVLYKVN.....QLKVIDNQRELRLSRELEP 338

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1 seqs, 1077 residues
Total number of hits satisfying chosen parameters: 1

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1 summaries

Database: 474982:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	283	16.1	1077	1 474982	ACCESSION:474982

ALIGNMENTS

RESULT 1	LOCUS	DEFINITION	ACCESSION	VERSION	DBSOURCE	KEYWORDS	SOURCE	ORGANISM	REFERENCE	AUTHORS	JOURNAL	COMMENT
474982	474982	C3G protein.	474982	GI:474982	locus HDMC3GP	accession D21239	Unknown.	Unknown.	1 (residues 1 to 1077)	Tanaka,S.	Unpublished (1994)	Submitted (20-Oct-1993) to DDBJ by: Michiyuki Matsuda

1-23-1 Toyama, Shinjuku-ku
Tokyo 162

Japan
Phone: 03-5285-1111 x2625
Fax: 03-5285-1150.

FEATURES
source
Protein

CD5
/product="C3G protein"
/function="ras guanine nucleotide releasing factor"
1..1077

/coded_by="D21239:131..3364"

Query Match 16.1%; Score 283; DB 1; Length 1077;
Best Local Similarity 28.3%; Pred. No. 0;
Matches 79; Conservative 48; Mismatches 98; Indels 54; Gaps 8;

QY	25	AIGLOP-DARGVATSLGLNERLFVNPEQVEHELIPHPDQGPYVGSAGGLDVSANDLAG 83
DB	818	ATSSQPLAARGVA-----ARPGTLHDFHS-----EIAE 846
QY	84	QLTDHMSLFNSIHQVELHYVLGPOHLRDVVTANLEFRMRRENELOYWATELCLCPVP 143
DB	847	QLTLDAELF---YKIRIPVLLMAKQONEKSPNLTQFTEHFNNMSYVWRSIIMQEA 903
QY	144	GPAQLRFRFKLAHLKEQKNINSFPAVWFGLSNSAISRLATHTWERLPHKVKRLYSALE 203
DB	904	QDRELLLKFKIKIMKHLKLNFNSTYALISALDSAPIRRL--EMOKQTSGLAEYCT-- 959
QY	204	RLDPSWNRVYRLALAKLSPPVIFPMPLLLKDMTFIHGNTLVENLINFER----- 256
DB	960	-LIDSSSFRAVRAALSEVEPCPIYIGLIDLTIFVHLGNPDYIDCKVNFSKRMQOFNI 10
QY	257	---MRMARAARMLHCRSHNPVPLSPLRVSRLHEDS 292
DB	1019	LDSWRCFOQA---HYDMRRNDIINFNFSDHLAEEA 1053

Search completed: October 8, 2003, 15:50:09
Job time: 0.001 secs

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OM protein - protein search, using SW model

Run on: October 8, 2003, 08:57:20 ; Search time 0.001 Seconds
(without alignments)
298.792 Million cell updates/sec

Title: us-09-856-679-2

Perfect score: 1759
Sequence: 1 MALAODGWTGQVLVKVN.....QLKVIDNQRELRLSRELEP 338

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 1 segs, 884 residues

Total number of hits satisfying chosen parameters: 1

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1 summaries

Database: 4079657.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1664	94.6	884	1	AAD12739 ACCESSION: AAD12739

ALIGNMENTS

RESULT 1
AAD12739 884 aa linear ROD 05-FEB-1999
LOCUS CAMP-regulated guanine nucleotide exchange factor 1 [Rattus norvegicus].
DEFINITION
ACCESSION AAD12739
VERSION AAD12739.1 GI:4079657
DBSOURCE locus RNU78167 accession U78167.1
KEYWORDS
SOURCE Rattus norvegicus (Norway rat)
ORGANISM Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
REFERENCE 1 (residues 1 to 884)
AUTHORS Kawasaki, H., Springle, G.M., Mochizuki, N., Toki, S., Nakaya, M., Matsuda, M., Housman, D.E. and Graybiel, A.M.
TITLE A family of CAMP-binding proteins that directly activate Rap1
JOURNAL Science 282 (5397), 2275-2279 (1998)
MEDLINE 99074384
PUBMED 9856955
REFERENCE 2 (residues 1 to 884)
AUTHORS Kawasaki, H., Housman, D.E. and Graybiel, A.M.
TITLE Direct Submission
JOURNAL Submitted (14-NOV-1996) Center for Cancer Research/Department of Brain and Cognitive Sciences, Massachusetts Institute of Technology, 77 Massachusetts Avenue, E17-540, Cambridge, MA 02135.

USA
FEATURES
source

Protein

CDS

Location/Qualifiers
1..884
/organism="Rattus norvegicus"
/db_xref="taxon:10116"
1..884
/product="CAMP-regulated guanine nucleotide exchange factor 1"
1..884
/gene="CAMP-GEF1"
/coded-by="U78167.1:197..2851"
/note="GEP domain shows highest similarity to RasGEF family; contains CAMP binding domain"

Query Match	Best Local Similarity	Score	DB 1:	Length	884:
Matches 316;	Conservative 11;	Mismatches 11;	Indels 0;	Gaps 0;	
QY	1	MALAODGWTGQVLVKVNSAGDAIGLPDARGVATSLGLNRLFPVNPQVHELIPHP	60		
DB	547	MAALAHEDHWTGQVLVKVNSAGDVVGLQPDARGVATSLGLNRLIFVVDPEVHELIPHP	60		
QY	61	DOLGPTVGSAGEGLDYSAKDLAQDLDDHDSLNSTHQVLLHYVGLPQHLRDVTYTNLE	120		
DB	607	EQLGPTLGSSEMLDLVSADKDLAQDLTEHDNLFNRIHQVELIHYVGLPQHLRDVTYTNLE	666		
QY	121	RFMRRENELOYVWATELCLCPVPGPRAQLRKFITKLAHLKEOKNLNLFPAVMEGLNSNA	180		
DB	667	RFMRRENELOYWATELCLCPVPGPRAQLRKFITKLAHLKEOKNLNLFPAVMEGLNSNA	726		
QY	181	ISRLAHTWERLPHKVRKLYSALERLLDPSVNNHRYRLALAKLSPPVTFEAPLLKDTFTI	240		
DB	727	ISRLAHTWERLPHKVRKLYSALERLLDPSVNNHRYRLALAKLSPPVTFEAPLLKDTFTI	786		
QY	241	HEGNHTLVENLIPEKRMARARARMLHRRSRINPPLSLRSRVSGLHSDQVARIKSTC	300		
DB	787	HEGNHTLVENLIPEKRMARARARMLHRRSRINPPLSLRSRVSGLHSDQVARIKSTC	846		
QY	301	SEQSLSTRSPASTWAVYQQLKVIDNQRELRLSRELEP	338		
DB	847	SEQSLSTRSPASTWAVYQQLKVIDNQRELRLSRELEP	884		

Search completed: October 8, 2003, 08:57:21
Job time: 1 sec